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≋ NCBI ●●● ● ● ⌐ ⌐ °° ●● ●●●●   ⌐Protein

PubMed   Nucleotide   Protein   Genome   Structure   PopSet

Search [Protein ▼] for [   ] [Go]

[Clear]

┌ Limits   Preview/Index   History   Clipboard

Display [GenPept ▼] [Save] [Text]   [Add to Clipboard]

┌ 1 : O65607 . DNA MISMATCH REPAI...[gi:3914062]   Related Sequences

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LOCUS        MSH3_ARATH    1076 aa                    PLN      15-DEC-1998
DEFINITION   DNA MISMATCH REPAIR PROTEIN MSH3.
ACCESSION    O65607
PID          g3914062
VERSION      O65607  GI:3914062
DBSOURCE     swissprot: locus MSH3_ARATH, accession O65607;
             class: standard.
             created: Dec 15, 1998.
             sequence updated: Dec 15, 1998.
             annotation updated: Dec 15, 1998.
             xrefs: gi: 5596409, gi: 2980796
             xrefs (non-sequence databases): PFAM PF00488
KEYWORDS     DNA repair; ATP-binding; DNA-binding.
SOURCE       thale cress.
  ORGANISM   Arabidopsis thaliana
             Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophy
             euphyllophytes; Spermatophyta; Magnoliophyta; eudicotyledons; c
             eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae;
             Arabidopsis.
REFERENCE    1   (residues 1 to 1076)
  AUTHORS    BEVAN,M., BARGUES,M., PEREZ-PEREZ,A., TEROL,J., TORRES,A.,
             PEREZ-ALONSO,M., HOHEISEL,J., MEWES,H.-W., MAYER,K. and
             SCHUELLER,C.
  TITLE      Direct Submission
  JOURNAL    Submitted (??-MAR-1998) to the EMBL/GenBank/DDBJ databases
  REMARK     SEQUENCE FROM N.A.
             STRAIN=CV. COLUMBIA
COMMENT      -------------------------------------------------------------
             This SWISS-PROT entry is copyright. It is produced through a
             collaboration between the Swiss Institute of Bioinformatics and
             the EMBL outstation - the European Bioinformatics Institute.
             The original entry is available from http://www.expasy.ch/sprot
             and http://www.ebi.ac.uk/sprot
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             [FUNCTION] NOT KNOWN. PROBABLE DNA-REPAIR PROTEIN.
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ORIGIN
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Revised: January 10, 2000.

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ID   ATM7J2      standard; DNA; PLN; 80386 BP.
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AC   AL022197;
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SV   AL022197.2
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DT   18-MAR-1998 (Rel. 55, Created)
DT   23-JUL-1999 (Rel. 60, Last updated, Version 5)
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DE   Arabidopsis thaliana DNA chromosome 4, P1 clone M7J2 (ESSA project)
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OS   Arabidopsis thaliana (thale cress)
OC   Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
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OC   core eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae;
OC   Arabidopsis.
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RN   [1]
RA   Bevan M., Bargues M., Perez-Perez A., Terol J., Torres A., Perez-Alonso M.,
RA   Hoheisel J., Mewes H.W., Mayer K.F.X., Lemcke K., Schueller C.;
RT   ;
RL   Unpublished.
XX
RN   [2]
RP   1-80386
RA   EU Arabidopsis sequencing project;
RT   ;
RL   Submitted (23-JUL-1999) to the EMBL/GenBank/DDBJ databases.
RL   MIPS, at the Max-Planck-Institut fuer Biochemie, Am Klopferspitz 18a,
RL   D-82152 Martinsried, FRG, E-mail:
RL   schuelle@mips.biochem.mpg.de,mayer@mips.biochem.mpg.de Project Coordinator:
RL   Mike Bevan, Molecular Genetics Department, Cambridge Laboratory, John Innes
RL   Centre, Colney Lane, NR4 7UJ Norwich, UK, E-mail: michael.bevan@bbsrc.ac.uk
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DR   SWISS-PROT; O65607; MSH3_ARATH.
DR   SWISS-PROT; P92965; RS40_ARATH.
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CC   Information on performance of analysis and a more detailed annotation
CC   of this entry and other sequences of chromosomes 3, 4 and 5 can be
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